

# SEQUENCE LISTING

<110> Ruben et al.

<120> Human Tumor Necrosis Factor Receptor TR17

<130> PF524P1

<140> Unassigned

<141> 2001-09-25

<150> 60/254,874

<151> 2000-12-13

<150> 60/235,991

<151> 2000-09-26

<150> 09/533,822

<151> 2000-03-24

<150> 60/188,208

<151> 2000-03-10

<160> 7

<170> PatentIn Ver. 2.1

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<213> Homo sapiens

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cag	gag	gag	cgc	ttt	cca	cag	ggc	ctg	tgg	acg	ggg	gtg	gct	atg	aga	96
Gln	Glu	Glu	Arg	Phe	Pro	Gln	Gly	Leu	Trp	Thr	Gly	Val	Ala	Met	Arg	
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tcc	tgc	ccc	gaa	gag	cag	tac	tgg	gat	cct	ctg	ctg	ggt	acc	tgc	atg	144
Ser	Cys	Pro	Glu	Glu	Gln	Tyr	Trp	Asp	Pro	Leu	Leu	Gly	Thr	Cys	Met	
		35					40					45				

tcc	tgc	aaa	acc	att	tgc	aac	cat	cag	agc	cag	cgc	acc	tgt	gca	gcc	192
Ser	Cys	Lys	Thr	Ile	Cys	Asn	His	Gln	Ser	Gln	Arg	Thr	Cys	Ala	Ala	
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ttc	tgc	agg	tca	ctc	agc	tgc	cgc	aag	gag	caa	ggc	aag	ttc	tat	gac	240
Phe	Cys	Arg	Ser	Leu	Ser	Cys	Arg	Lys	Glu	Gln	Gly	Lys	Phe	Tyr	Asp	
65					70					75					80	

cat	ctc	ctg	agg	gac	tgc	atc	agc	tgt	gcc	tcc	atc	tgt	gga	cag	cac	288
His	Leu	Leu	Arg	Asp	Cys	Ile	Ser	Cys	Ala	Ser	Ile	Cys	Gly	Gln	His	
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Pro Lys Gln Cys Ala Tyr Phe Cys Glu Asn Lys Leu Arg Ser Pro Val	
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aac ctt cca cca gag ctc agg aga cag cgg agt gga gaa gtt gaa aac	384
Asn Leu Pro Pro Glu Leu Arg Arg Gln Arg Ser Gly Glu Val Glu Asn	
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aat tca gac aac tcg gga agg tac caa gga ttg gag cac aga ggc tca	432
Asn Ser Asp Asn Ser Gly Arg Tyr Gln Gly Leu Glu His Arg Gly Ser	
130 135 140	
gaa gca agt cca gct ctc ccg ggg ctg aag ctg agt gca gat cag gtg	480
Glu Ala Ser Pro Ala Leu Pro Gly Leu Lys Leu Ser Ala Asp Gln Val	
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Ala Leu Val Tyr Ser Thr Leu Gly Leu Cys Leu Cys Ala Val Leu Cys	
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tgc ttc ctg gtg gcg gtg gcc tgc ttc ctc aag aag agg ggg gat ccc	576
Cys Phe Leu Val Ala Val Ala Cys Phe Leu Lys Lys Arg Gly Asp Pro	
180 185 190	
tgc tcc tgc cag ccc cgc tca agg ccc cgt caa agt ccg gcc aag tct	624
Cys Ser Cys Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser	
195 200 205	
tcc cag gat cac gcg atg gaa gcc ggc agc cct gtg agc aca tcc ccc	672
Ser Gln Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro	
210 215 220	
gag cca gtg gag acc tgc agc ttc tgc ttc cct gag tgc agg gcg ccc	720
Glu Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro	
225 230 235 240	
acg cag gag agc gca gtc acg cct ggg acc ccc gac ccc act tgt gct	768
Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys Ala	
245 250 255	
gga agg tgg ggg tgc cac acc agg acc aca gtc ctg cag cct tgc cca	816
Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro Cys Pro	
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cac atc cca gac agt ggc ctt ggc att gtg tgt gtg cct gcc cag gag	864
His Ile Pro Asp Ser Gly Leu Gly Ile Val Cys Val Pro Ala Gln Glu	
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<212> PRT

<213> Homo sapiens

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 His Leu Leu Arg Asp Cys Ile Ser Cys Ala Ser Ile Cys Gly Gln His  
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 Pro Lys Gln Cys Ala Tyr Phe Cys Glu Asn Lys Leu Arg Ser Pro Val  
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 115 120 125  
 Asn Ser Asp Asn Ser Gly Arg Tyr Gln Gly Leu Glu His Arg Gly Ser  
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 145 150 155 160  
 Ala Leu Val Tyr Ser Thr Leu Gly Leu Cys Leu Cys Ala Val Leu Cys  
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 Cys Phe Leu Val Ala Val Ala Cys Phe Leu Lys Lys Arg Gly Asp Pro  
 180 185 190  
 Cys Ser Cys Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser  
 195 200 205  
 Ser Gln Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro  
 210 215 220  
 Glu Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro  
 225 230 235 240  
 Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys Ala  
 245 250 255  
 Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro Cys Pro  
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 tctcccggac tcttgaggtc acatgcgtgg tgggtggacgt aagccacgaa gaccctgagg 180  
 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg 240  
 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact 300  
 ggctgaatgg caaggagtac aagtgcgaagg tctccaacaa agccctccca acccccatcg 360  
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 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctgggc aaaggcttct 480  
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 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc 660  
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